



## Data Article

# Draft genome sequence of *Brevibacterium epidermidis* EZ-K02 isolated from nitrocellulose-contaminated wastewater environments



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## ABSTRACT

*Brevibacterium* spp. are aerobic, nonbranched, asporogenous, gram-positive, rod-shaped bacteria which may exhibit a rod-coccus cycle when cells get older and can be found in various environments. Several *Brevibacterium* species have industrial importance and are capable of biotransformation of various contaminants. Here we describe the draft genome sequence of *Brevibacterium epidermidis* EZ-K02 isolated from nitrocellulose-contaminated wastewater environments. The genome comprises 3,885,924 bp, with a G + C content of 64.2%. This whole genome shotgun project has been deposited at DDBJ/ENA/GenBank under the accession PDHL00000000.

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